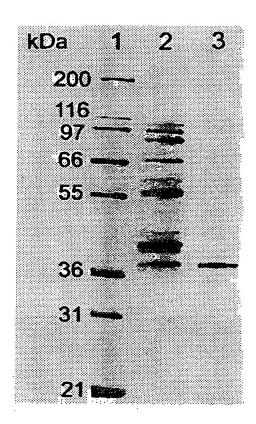
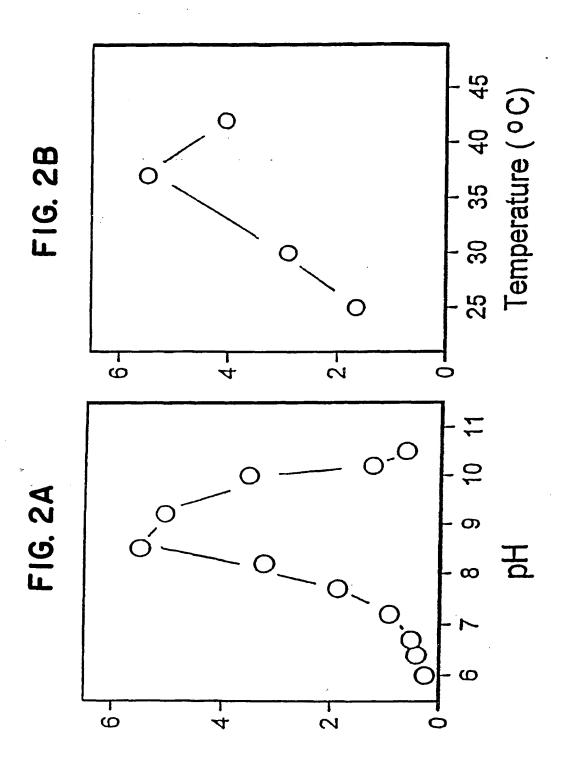
1/8

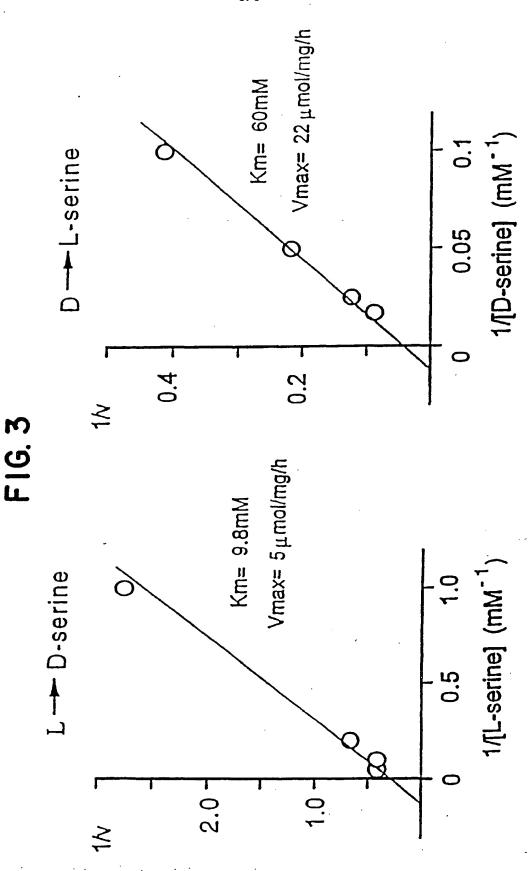
FIG. I





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0988260091889809

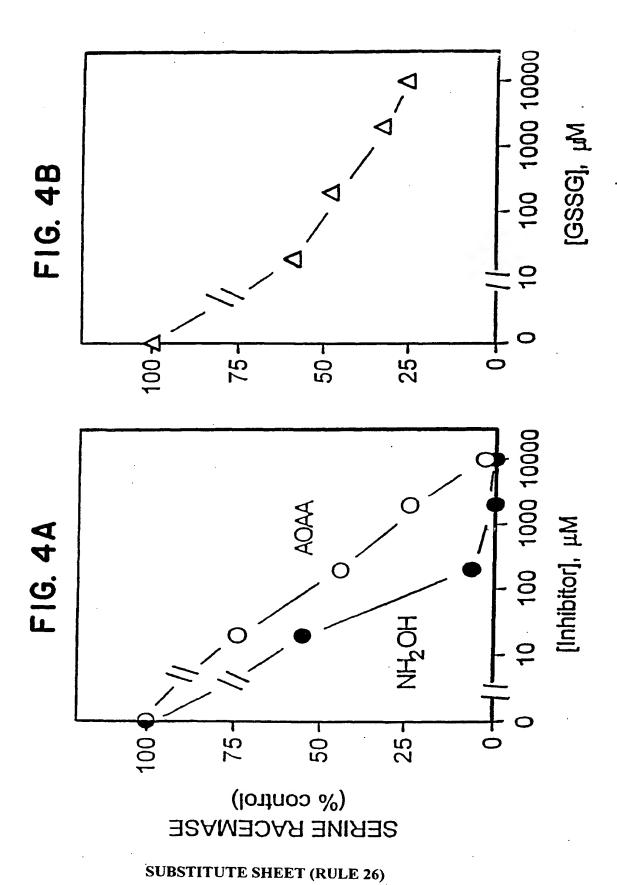
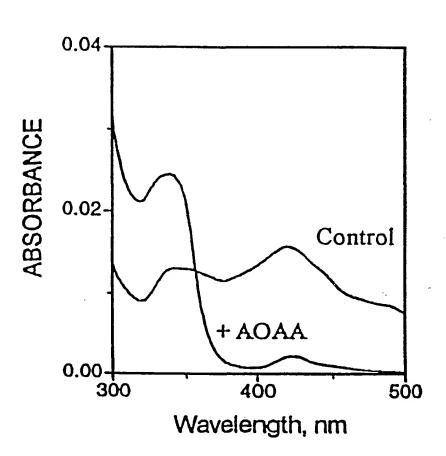


FIG. 5



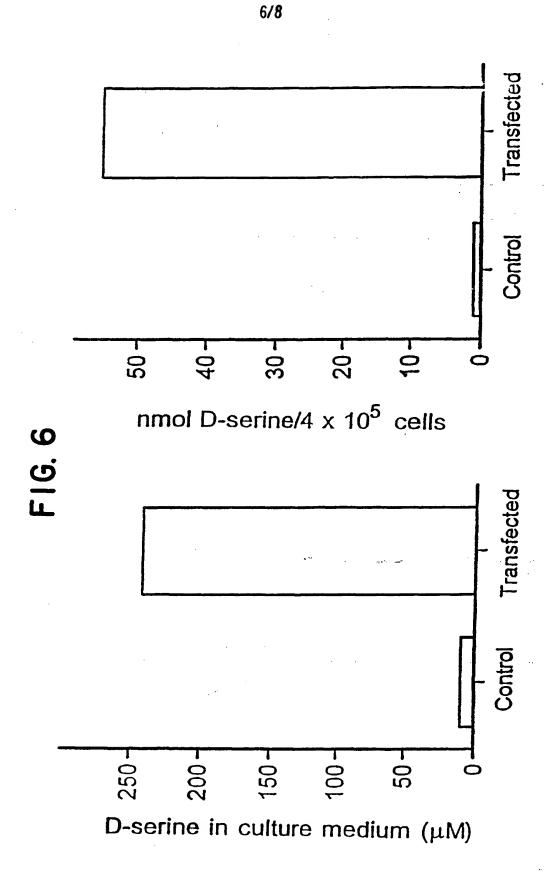


FIG. 7A

ads complete gene, racemase serine musculus

Source: house mouse

Organism : Mus musculus

Eutheria; Mammalia; Eukaryota; Metazoa; Chordata; Vertebrata;

Rodentia; Sciurognathi; Muridae; Murinae; Mus

Strain = "Balb/c"

Dev. Stage = Adult

Tissue type = brain

Gene Starts at 219 and ends 1238

AGTCCTTACC GAAACAGAAG ACAATTGCCC CCATAAAATG TGGTCTGCTG CTCACCCCAG AAATGTGAGC AGAGGCTTAA GGAAACCATG GTGGTTCCCC ATAGTATACT TACTGCATCT GGCCAATCAA TGGACAAGGA GTGTGCTCAG CTCTATCCAC TGGAGCATCG AATTATGCAA AGTATGGAAG TAATGCCATC TCACAGCAGC TGCTTACATT TCTTTTCTTC CGGAGCCAGC GGCCAAGATG ATCCCTCTTC TCCTGCAGTA ACATTCAAGA CAGGGGGCAA TCCAAGCCTA TCACTCAAAG CAGTGATAGC AGAGAACCAT AAGGAATTCC GAGGTGCCCT CCGTAGTTAC ACTGGTCCTG TTGAGAGAGA CAGCCATGTG CAGAGGCAGC AATCAAATAG GCTAAACTGG CAGGAGCCTG GCTCATATICA AGAGAAAAGG TTTAAGATTC AAGCCCAAAG AAACTGGCAA CCTTTGCCAC CTCCATTTTG CAACTGCAAG TGACGAGTCC CCATCCCAAC AACTGGGTCT GCCAGAAGAG TGAACTGAAA GAGCTGTCAG CACCTATGCT GCCGCCGTGC TGTTGAAAAA GACCTTACAC TAAAACACAG TGAGAAGGAA CCTTTGCTGA TGCTAACAAG TCTTCCAGAA GTGACCCAAG GCATCTTGGT CAGAATTTTT PTCCTGACAC AAACAGCTCC GCCAAGCTCT 181 361 541 601 241 301 421 481

8/8

GGAGGAGGAG AAGGTATACG GAACTGACCC GGCTTGAATA GATGAAATCA ACTGCTGGCG ATGAAGAAG GCAGTAGAGA CANANCTICCA GTAAAGAACG GTGGGGCAGG TTGTCTCTAG ATACAGAGCA GAGATGTCAC GGAAATACTC CCCTTGGTAG ATGCACTGGT GGTACCAGTA ACCTAGTGTG ACTGAAAGGA ATCCAGCATT CATTGAGCCG CCTGAACTGG GACAACTGAA TTGGTTGGTT CTCTCCAGAA TCAGGCAAGA CAAAATCCTA AGTCCACTTG CTTTCCCCAT TTGGTCAAAA TGTCACCGAA CACTGTTGAA ACCAGTCTAA ATGGTGTCAA ATGTCTTCAC AAACAGATAA AGGCCCTGAA GAGATTTCAG TGAAACTGCT TCCAAACAGT CTGTTTAAAT CAGAGACATG CACTGGAGTT CTACTTCAGA ATGGGTCATA ACCTAACCTC AAATTATGTT CTAAAAGTTA ATTACAATTA CTTGTGGATG CAGACTCCAA GATGACTGCT ACCATAGCAG TGGGGGAGAA TCTCAGCATT GGGAATGTAG CAGACGGTTT TYCCCTYTAA CTGTACTTAG TACTCCATGT TCTGGTACAT ACTGTGTCCT GAACCAGGTT TGGAATAGCC ATTCATTAAT CTCGAATGCA TCCTCCAGAA CCAGCTGGTG CCATGGGTAC **FATAAGAGAC** TGCAGTGCTG TGTTTCCATC CATGTAAATT ACTCAGTGGG AGCTCCTTAC GGTCACAACC CTGAGTCTCT AGGTACTGGC TGGAAGTGCT GAATGGTTGC CTGCTGAGCC CCAATCTTCA CCTGGCCTAT AGTATGCAAC TGGCACTGGC TCTGCATTGT CTGAACGGCC CAGAACCCAA TTTTAACCCA GCTTATAGAG CTTTATATGG ATGAAAATTT TGTAAGTAAT CAAGCTGCCT 841 901 961 1021 1081 1201 1261 1381 1561 1621 1141 1321 1441 1501

Translation:="MCAQYCISFADVEKAHINIQDSIHLTPVLTSSILNQIAGRNLFFKCELFQKTGSFKIR galnairglipdtpeekpkavvthssgnhgqaltyaaklegipayivvpqtapnckklaiqaygasivycdp SDESREKVTQRIMQETEGILVHPNQEPAVIAGQGTIALEVLNQVPLVDALVVPVGGGGMVAGIAITIKALKP SVKVYAAEPSNADDCYQSKLKGELTPNLHPPETIADGVKSSIGLNTWPIIRDLVDDVFTVTEDEIKYATQLV WGRMKLLIEPTAGVALAAVLSQHFQTVSPEVKNVCIVLSGGNVDLTSLNWVGQAERPAPYQTVSV~